

Figure 1B

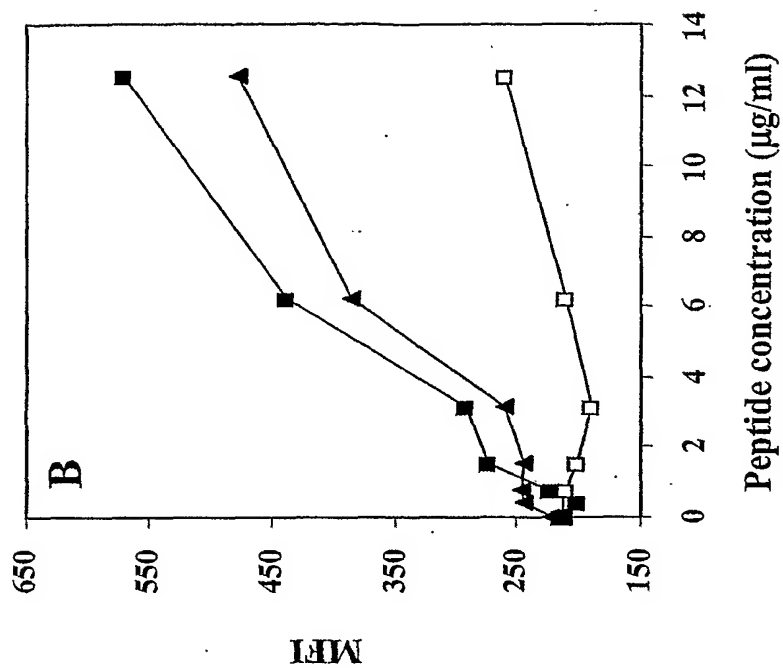
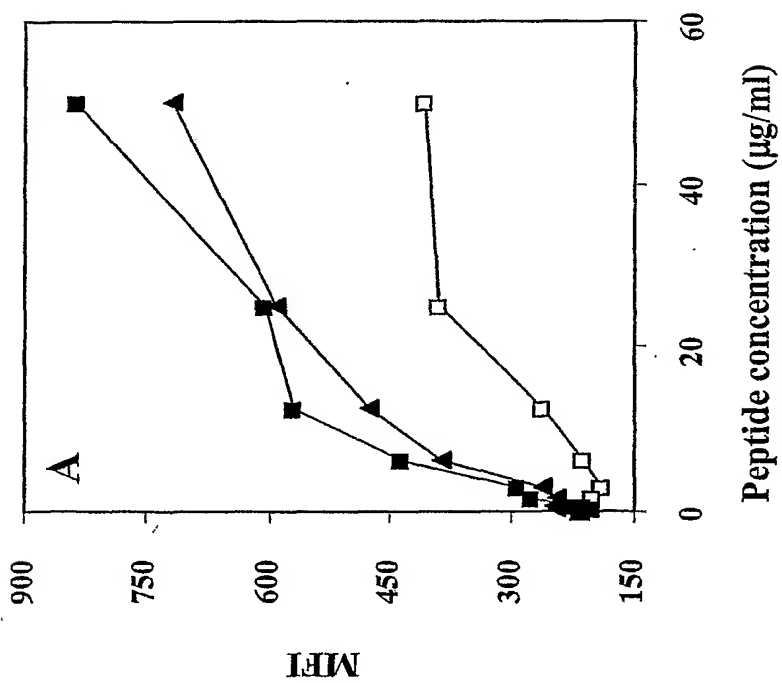
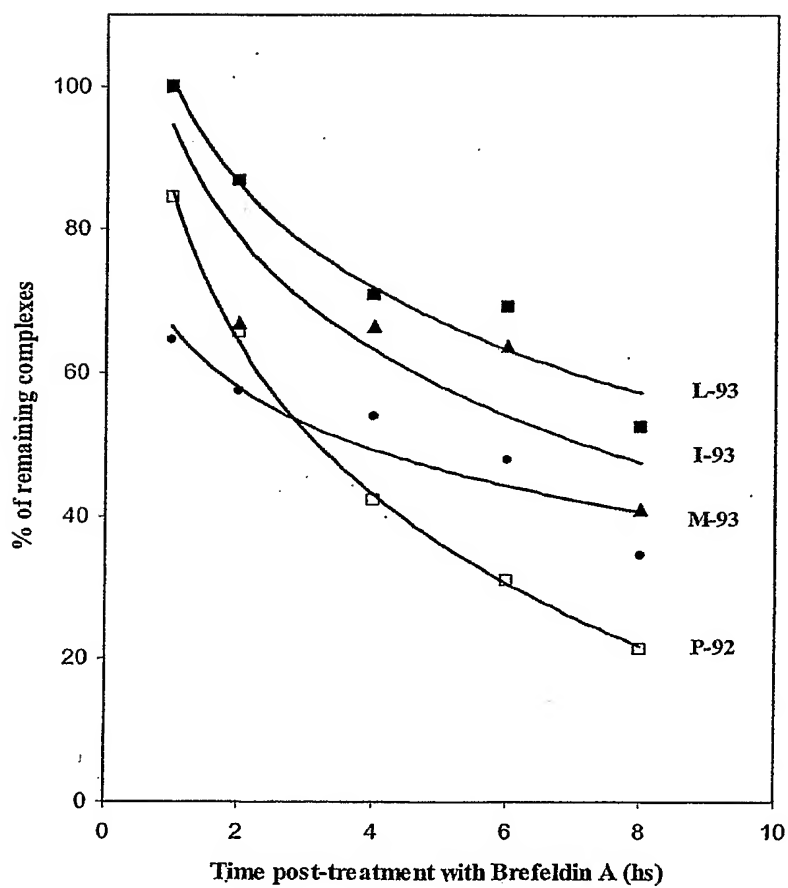


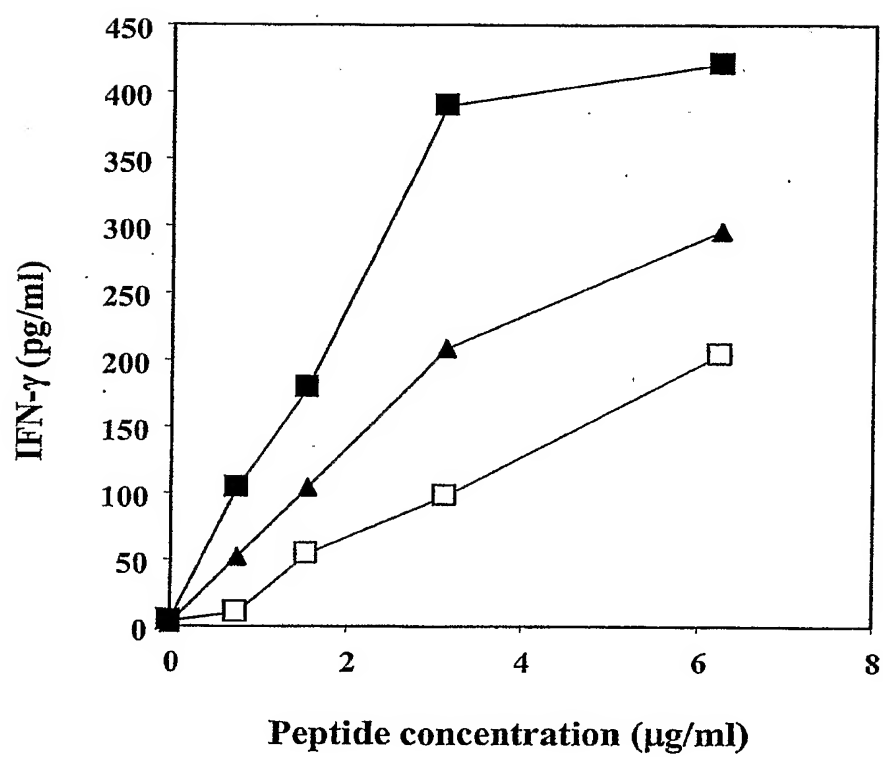
Figure 1A



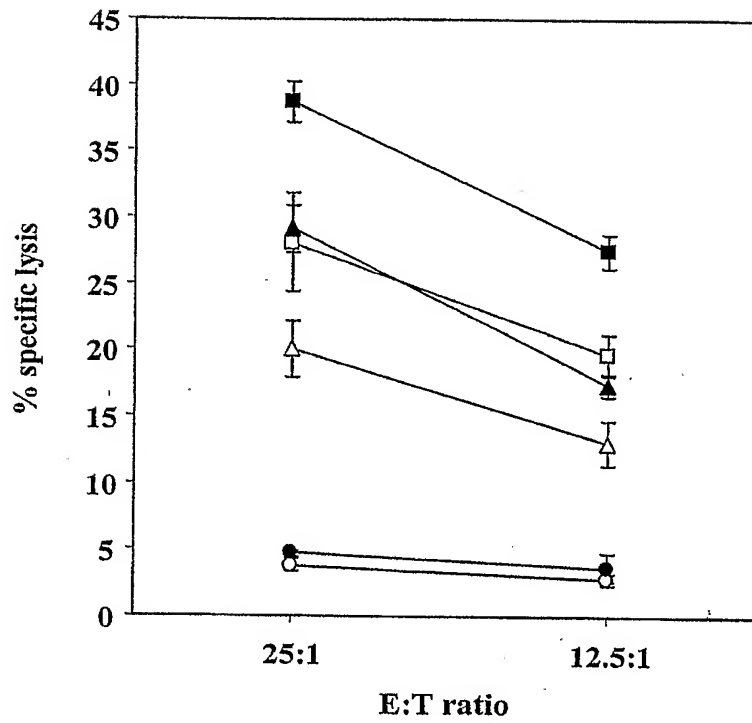
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Figure 2

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Figure 3

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Figure 4

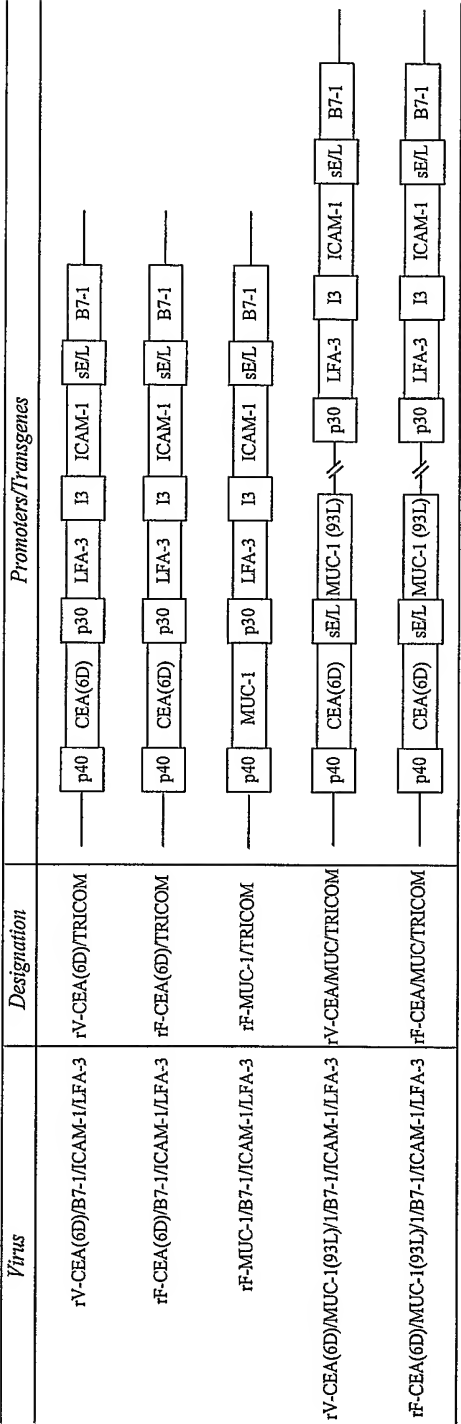
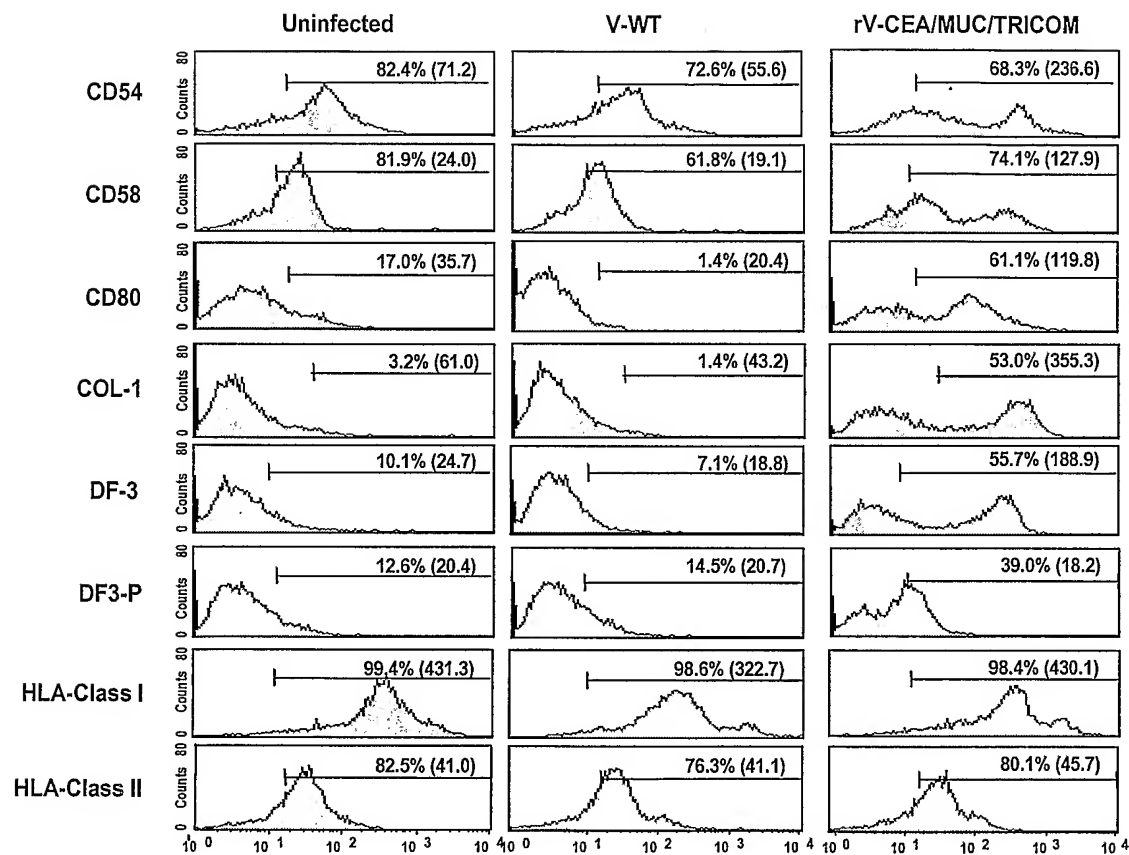
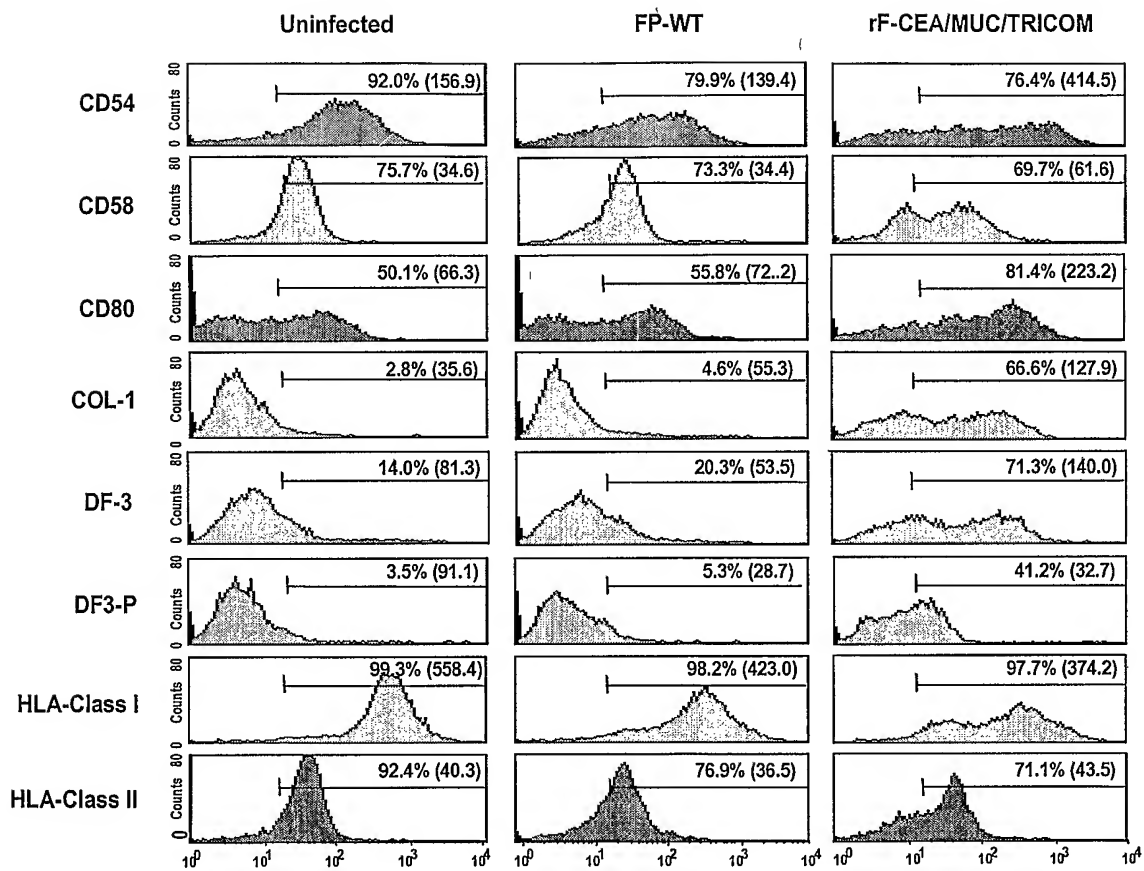


Figure 5

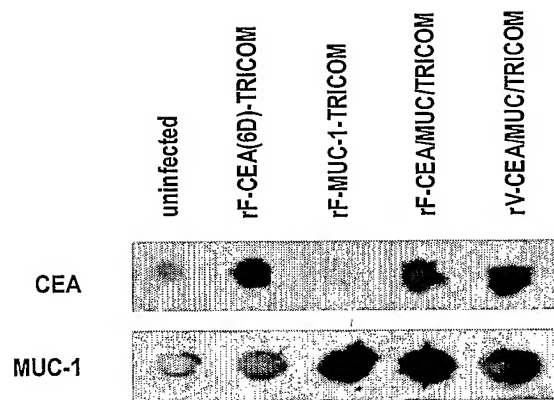
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**Figure 6**

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**Figure 7**

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**Figure 8**

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1  ATGACACCGG  GCACCCAGTC  TCCTTCTTTC  CTGCTGCTGC  TCCTCACAGT  GCTTACAGTT
61  GTTACGGGTT  CTGTCTATGC  AAGCTCTACC  CCAGGTGGAG  AAAGGAGAC  TTCGGCTAAC
121  CAGAGAGATT  CAGTGGCCAG  CTCTACTGAG  AGAATCTG  TGRGTATGAC  TAGCTCCGTA
181  CTCTCCAGGC  ACAGCCCGG  TTCAGGCTCC  TCCACCACTC  AGGACACGGA  TGTCACTCTG
241  GCCCCGGCCA  CGGACACG  CTGAGCTTCA  GCTGGCTTGT  GGGACACGGA  TGTCACTCTG
301  GTACCAAGTA  CTAGACACG  TTAGGTAGC  ACAGCACTC  CTGCTCATGG  AGTAACTAGT
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781  ACTCTTACCA  CCTCTGCCG  CCATAGCAC  AGACTGATG  CCAGTAGCAC  TCACTCTAGC
841  ACGGTACCTC  CTCTCACCCT  CTCCATCAC  AGCACTCTC  CCGAGTGTG  TACTGGGTC
901  TCTTCTTTT  TCTGTCTTT  TCACATTTCA  AACCTCCAGT  TTAATTCCT  TCTGGAGAT
961  CCCAGCACCG  ACTACTACCA  AGAGCTGCAG  AGAGACATTT  CTGAATGTT  TTTSCAGATT
1021  TATAACACAG  GGGTTTCT  GGGCTCTCC  AATATTAGT  TCCAGCCAG  ATCTGTGGT
1081  GTACATTTGA  CTCTGGCTT  CCGAGAGGT  ACCATCAATG  TCCAGCCAGT  AGAGCTCAGC
1141  TTCAATCAGT  ATAAACGGA  AGCAGCTCT  CGATTTAC  TGACCATCTC  AGAGCTCAGC
1201  GTGAGTATG  TSCCATTTCC  TTCTCTGCC  CAGTCTGGG  CTGGGTGCC  AGCTGGGGC
1261  ATCGGCTGC  TGGTCTGCT  CTGTCTCTG  GTTGGGTCG  TGGATCTCT  TCCAGCCCGG
1321  TTGGCTGTCT  GTCTGTGCG  CCGAAGAAC  TAGGGCAGC  TGGATCTCT  TCCAGCCCGG
1381  GATACCTACC  ATCTATGAG  CGAGTACCC  CCTACACCA  CCCATGGCG  CTATGTGCC
1441  CCTAGCAGTA  CCGATCGTAG  CCCCTATGAG  AGGTTTCTG  CAGGTAATG  TGGCAGCAGC
1501  CTCTCTTACA  CAAACCCAGC  AGTGGCAGC  ACTTCTGCCA  ACTTGTAG

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Figure 9

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Figure 10

MTPGTQSPFFLLLLTLTVVTGSGHASSTPGGEKETSATQSSVPSSTEKNAV
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